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RAW SEQUENCE LISTING
PATENT APPLICATION: US/09/846,091

DATE: 03/19/2002
TIME: 14:53:33

Input Set : N:\Crf3\RULE60\09846091.raw
Output Set: N:\CRF3\03192002\I846091.raw

ENTERED

<110> APPLICANT: HAYNES, Joel R.
MACKLIN, Michael D.
PAYNE, Lendon G.
<120> TITLE OF INVENTION: NUCLEIC ACID IMMUNIZATION
<130> FILE REFERENCE: APF40
<140> CURRENT APPLICATION NUMBER: 09/846,091
<141> CURRENT FILING DATE: 2001-04-30
<150> PRIOR APPLICATION NUMBER: US/09/561,951
<151> PRIOR FILING DATE: 2000-05-01
<160> NUMBER OF SEQ ID NOS: 11
<170> SOFTWARE: PatentIn Ver. 2.1
<210> SEQ ID NO: 1
<211> LENGTH: 24
<212> TYPE: PRT
<213> ORGANISM: Influenza A virus
<400> SEQUENCE: 1
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Cys Arg Cys Asn Gly Ser Ser Asp
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<212> TYPE: PRT
<213> ORGANISM: Influenza A virus
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Cys Arg Cys Asn Asp Ser Ser Asp
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<210> SEQ ID NO: 3
<211> LENGTH: 24
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<213> ORGANISM: Influenza A virus
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<210> SEQ ID NO: 4
<211> LENGTH: 20
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<213> ORGANISM: Artificial Sequence
<220> FEATURE:

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<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic Construct

<400> SEQUENCE: 4
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<210> SEQ ID NO: 5
<211> LENGTH: 20
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<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic Construct

<400> SEQUENCE: 5
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<211> LENGTH: 65
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<213> ORGANISM: Artificial Sequence
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<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic Construct

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<211> LENGTH: 28
<212> TYPE: DNA
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic Construct

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<210> SEQ ID NO: 8
<211> LENGTH: 17
<212> TYPE: PRT
<213> ORGANISM: Artificial Sequence
<220> FEATURE:
<223> OTHER INFORMATION: Description of Artificial Sequence: Synthetic Construct

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1 5 10 15
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<210> SEQ ID NO: 9
<211> LENGTH: 294
<212> TYPE: DNA
<213> ORGANISM: Influenza A/Kagoshima/10/95(H3N2)
<220> FEATURE:
<221> NAME/KEY: CDS

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109         1               5                  10                  15
110       tgc aga tgc aac ggt tca agt gac ccg ctt gtt gtt gct gcg agt atc   96
111       Cys Arg Cys Asn Gly Ser Ser Asp Pro Leu Val Val Ala Ala Ser Ile
112                    20                  25                  30
113       att ggg atc ttg cac ttg ata ttg tgg att ttt gat cgt ctt ttt ttc  144
114       Ile Gly Ile Leu His Leu Ile Leu Trp Ile Phe Asp Arg Leu Phe Phe
115                35                  40                  45
116       aaa tgc atc tat cga ctc ttc aaa tac ggt ctg aaa aga ggg cct tct  192
117       Lys Cys Ile Tyr Arg Leu Phe Lys Tyr Gly Leu Lys Arg Gly Pro Ser
118            50                  55                  60
119       acg gaa gga gta cct gag tct atg agg gaa gaa tat cga aag gaa cag  240
120       Thr Glu Gly Val Pro Glu Ser Met Arg Glu Glu Tyr Arg Lys Glu Gln
121         65                  70                  75                  80
122       cag aat gct gtg gat gct gac gac agt cat ttt gtc agc ata gag ctg  288
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124                       85                  90                  95
125       gag taa                                                        294
126       Glu
128 <210> SEQ ID NO: 10
129 <211> LENGTH: 97
130 <212> TYPE: PRT
131 <213> ORGANISM: Influenza A/Kagoshima/10/95(H3N2)
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134         1               5                  10                  15
135       Cys Arg Cys Asn Gly Ser Ser Asp Pro Leu Val Val Ala Ala Ser Ile
136                    20                  25                  30
137       Ile Gly Ile Leu His Leu Ile Leu Trp Ile Phe Asp Arg Leu Phe Phe
138                35                  40                  45
139       Lys Cys Ile Tyr Arg Leu Phe Lys Tyr Gly Leu Lys Arg Gly Pro Ser
140            50                  55                  60
141       Thr Glu Gly Val Pro Glu Ser Met Arg Glu Glu Tyr Arg Lys Glu Gln
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148 <211> LENGTH: 4622
149 <212> TYPE: DNA
150 <213> ORGANISM: pM2-FL plasmid
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154       taggtggacc agttggtgat tttgaacttt tgctttgcca cggaacggtc tgcgttgtcg 180
155       ggaagatgcg tgatctgatc cttcaactca gcaaaagttc gatttattca acaaagccgc 240
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156 cgtcccgtca agtcagcgta atgctctgcc agtgttacaa ccaattaacc aattctgatt 300
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159 ggatggcaag atcctggtat cggtctgcga ttccgactcg tccaacatca atacaaccta 480
160 ttaatttccc ctcgtcaaaa ataaggttat caagtgagaa atcaccatga gtgacgactg 540
161 aatccggtga gaatggcaaa agcttatgca tttctttcca gacttgttca acaggccagc 600
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163 cctgagcgag acgaaatacg cgatcgctgt taaaaggaca attacaaaca ggaatcgaat 720
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205 cattttgtca gcatagagct ggagtaagga tcctcgcaat ccctaggagg attaggcaag 3240
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